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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
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EXAMINER

FREDMAN, JEFFREY NORMAN

ART UNIT	PAPER NUMBER
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1634

DATE MAILED: 12/04/2003

Please find below and/or attached an Office communication concerning this application or proceeding.

<b>Office Action Summary</b>	<b>Application No.</b> 09/827,383	<b>Applicant(s)</b> MITTMANN ET AL.	
	<b>Examiner</b> Jeffrey Fredman	<b>Art Unit</b> 1634	

-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

**Period for Reply**

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If the period for reply specified above is less than thirty (30) days, a reply within the statutory minimum of thirty (30) days will be considered timely.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133).
- Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

**Status**

- 1) ☒ Responsive to communication(s) filed on 06 October 2003.
- 2a) ☐ This action is **FINAL**.                      2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

**Disposition of Claims**

- 4) ☒ Claim(s) 1-14 is/are pending in the application.
- 4a) Of the above claim(s) 13 and 14 is/are withdrawn from consideration.
- 5) ☐ Claim(s) \_\_\_\_\_ is/are allowed.
- 6) ☒ Claim(s) 1-12 is/are rejected.
- 7) ☐ Claim(s) \_\_\_\_\_ is/are objected to.
- 8) ☐ Claim(s) \_\_\_\_\_ are subject to restriction and/or election requirement.

**Application Papers**

- 9) ☐ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on \_\_\_\_\_ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.  
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).  
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

**Priority under 35 U.S.C. §§ 119 and 120**

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).  
a) ☐ All   b) ☐ Some \*   c) ☐ None of:  
1. ☐ Certified copies of the priority documents have been received.  
2. ☐ Certified copies of the priority documents have been received in Application No. \_\_\_\_\_.  
3. ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).
- \* See the attached detailed Office action for a list of the certified copies not received.
- 13) ☒ Acknowledgment is made of a claim for domestic priority under 35 U.S.C. § 119(e) (to a provisional application) since a specific reference was included in the first sentence of the specification or in an Application Data Sheet. 37 CFR 1.78.  
a) ☐ The translation of the foreign language provisional application has been received.
- 14) ☐ Acknowledgment is made of a claim for domestic priority under 35 U.S.C. §§ 120 and/or 121 since a specific reference was included in the first sentence of the specification or in an Application Data Sheet. 37 CFR 1.78.

**Attachment(s)**

- |   |   |
|---|---|
| 1) <input checked="" type="checkbox"/> Notice of References Cited (PTO-892)                             | 4) <input type="checkbox"/> Interview Summary (PTO-413) Paper No(s). _____  |
| 2) <input type="checkbox"/> Notice of Draftsperson's Patent Drawing Review (PTO-948)                    | 5) <input type="checkbox"/> Notice of Informal Patent Application (PTO-152) |
| 3) <input checked="" type="checkbox"/> Information Disclosure Statement(s) (PTO-1449) Paper No(s) _____ | 6) <input type="checkbox"/> Other: _____                                    |

## **DETAILED ACTION**

### ***Election/Restrictions***

1. Applicant's election without traverse of Group I, claims 1-10, and of SEQ ID NO:s 1-10 in the paper filed October 6, 2003, is acknowledged.

### ***Claim Rejections - 35 USC § 101***

2. 35 U.S.C. 101 reads as follows:

Whoever invents or discovers any new and useful process, machine, manufacture, or composition of matter, or any new and useful improvement thereof, may obtain a patent therefor, subject to the conditions and requirements of this title.

3. Claims 1-12 are rejected under 35 U.S.C. 101 because the claimed invention lacks patentable utility.

The current claims are drawn to a set of nucleic acids tags with at least 10 sequences whose sequences comprise SEQ ID NO: 1-2050, with SEQ ID Nos: 1-10 being selected.

### **Credible Utility**

Following the requirements of the Utility Guidelines (See: Federal Register: December 21, 1999 (Volume 64, Number 244), revised guidelines for Utility.), the first inquiry is whether a credible utility is cited in the specification for use of the nucleic acids. One cited utility identified in the specification is to analyze genomic DNA (see page 10, for example). This utility is credible.

Upon identification of credible utilities, the next issue is whether there are any well established utilities for the nucleic acid. No well established utilities for these specific SEQ ID Nos: 1-10 are identified in either the specification or in the cited prior art.

**Substantial utility**

Given the absence of a well established utility, the next issue is whether substantial utilities are disclosed in the specification. Here, there is no evidence of any substantial utility. No substantial use for a set of sequences comprising SEQ ID NOs: 1-10 is found in the specification nor is there any use for the method or system involving SEQ ID NO: 1.

As noted in the utility guidelines, methods of treating unspecified diseases, basic research on a product to identify properties, intermediate products which themselves lack substantial utility are all insubstantial utilities (see page 6 of the Utility guideline training materials). If there were evidence of the association of SEQ ID NO: 1 with any disease state, with a protein activity or with some other biological phenotype, this evidence might be considered regarding a substantial utility. However, no such evidence is found. At best, the utilities of analyzing genomic DNA are indicative that SEQ ID NOs: 1-10 are intermediate products which lack substantial utility.

In the Supreme Court case of *Brenner v. Manson*, 148 U.S.P.Q. 689 (1966), a novel compound which was structurally analogous to other compounds that were known to possess anti-tumor activity was alleged to be potentially useful as an anti-tumor agent in the absence of evidence supporting this utility. The court expressed the opinion that all chemical compounds are "useful" to the chemical arts when this term is given its broadest interpretation. However, the court held that this broad interpretation was not the intended definition of "useful" as it appears in 35 U.S.C. §101, which requires that

an invention must have either an immediately apparent or fully disclosed "real world" utility. The court held that:

The basic quid pro quo contemplated by the Constitution and the Congress for granting a patent monopoly is the benefit derived by the public from an invention with substantial utility. . . . [u]nless and until a process is refined and developed to this point-where specific benefit exists in currently available form-there is insufficient justification for permitting an applicant to engross what may prove to be a broad field. . . . a patent is not a hunting license. . . . [i]t is not a reward for the search, but compensation for its successful conclusion.

The instant claims are drawn to a set of polynucleotides with no known function. The specification does not teach the function of any of the nucleic acids to which these sequences hybridize. The function of these nucleic acids is as yet undetermined with no known biological significance. There is no evidence of record or any line of reasoning that would support a conclusion that the nucleic acid of the instant application was, as of the filing date, useful for any specific assay or therapeutic use. Until some actual and specific significance can be attributed to the nucleic acid, one of ordinary skill in the art would be required to perform additional experimentation in order to determine how to use the claimed invention. Following both the Utility Guidelines and the direction of the Supreme Court of the United States in *Brenner*, there is no specific benefit in using a set of these specific sequences. Thus, there is no immediately substantial or "real world" utility as of the filing date.

### **Specific Utility**

In the current case, even if the substantial utility argument above were found unpersuasive, there is clearly no specific utility. To the extent that the nucleic acid and polymorphisms in SEQ ID NO: 1-10 can be used in genomic analysis assays, this utility

is not at all specific to SEQ ID NO: 1-10. Literally any sequence would function in a genome analysis assay as described in the specification. As the utility guideline training materials note on page 5-6 regarding specific utility that "a claim to a polynucleotide whose use is disclosed simply as a 'gene probe' or 'chromosome marker' would not be considered to be *specific* in the absence of a disclosure of a specific DNA target (*italics in original*)". Here, there is no disclosure of any specific use of SEQ ID NO: 1-10 that is not shared with any other sequence.

Further, the sequences are not even species or chromosome specific, based upon the sequence search. As the attached search of SEQ ID NO: 3 in Genbank demonstrates, result 3 shows an 90% match (local similarity) to a sequence in chromosome 14 of humans while result 13 shows an 89.5% match (local similarity) to human chromosome 8. Further, the remaining results show similarity to *Pseudomonas*, Lotus, Rats and Rice with equivalent levels of local similarity. Similar results exist for the other 9 probes. So the sequences claimed lack 100% specificity to any particular organism in Genbank, and the specification lacks any discussion of the target for these oligonucleotides. Consequently, there is no specific target for any of the claimed sequences. With regard to the utility analysis, the current situation directly tracks Example 9 of the utility guidelines, where an unknown nucleic acid fragment of entirely unknown function was characterized as lacking utility.

Finally, there appears to be no element which is unique to the selected sequences. That is, the ability of the array to be used in SNP-IT™ assays, for example, is not sequence dependent. That is, there is nothing specific to the 2050 sequences of

the current claim which distinguish these sequences from a different set of 2050 sequences or from any set of 2050 unrelated sequences.

Therefore, a set of nucleic acids comprising SEQ ID NO: 1-10 has no specific utility.

***Claim Rejections - 35 USC § 112***

4. The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

5. Claims 1-12 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the enablement requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to enable one skilled in the art to which it pertains, or with which it is most nearly connected, to make and/or use the invention.

**Nature of Invention**

The current claims drawn to a set of nucleic acids tags with at least 10 sequences whose sequences comprise SEQ ID NO: 1-2050, with SEQ ID Nos: 1-10 being selected. The nature of this invention is in nucleic acid analysis of a particular sequence with no other associated information. The invention is in an class of invention which the CAFC has characterized as "the unpredictable arts such as chemistry and biology." *Mycogen Plant Sci., Inc. v. Monsanto Co.*, 243 F.3d 1316, 1330 (Fed. Cir. 2001).

**Breadth of the claims**

The claims are drawn to a set of nucleic acids comprising at least 10 sequences from SEQ ID Nos: 1-2050, and for which SEQ ID Nos: 1-10 were selected.

**Amount of Guidance in the Specification**

The specification discloses the entire sequence of SEQ ID NOs: 1-10, but identifies no particular use for the sequence. In particular, the specification lacks any discussion of the target of SEQ ID NOS: 1-10, or of any of the 2040 other sequences. As noted in the utility rejection above, this utility is not found to be substantial nor specific and consequently, the specification provides NO guidance regarding how to use SEQ ID NO: 1. The general guidance that the method is useful in methods of genomic analysis fails to provide the specific details necessary to apply sequences whose targets are unknown to such genomic analysis.

**Working Examples**

There are working examples in which the sequences are hybridized. However, there are no working examples in which SEQ ID NOs: 1-10, or indeed any of the 2040 other sequences, are used in any assay for detection or diagnosis of any disease or any other related utility. No real world use or particular use is given for these sequences.

**Amount of Guidance in Prior Art**

As noted in the utility rejection above, the prior art provides no guidance with regard to the particular function of SEQ ID NOs: 1-10.

**Skill in the Art**



While no evidence is adduced, the examiner believes the skill in the art would be considered high.

### **Predictability of the Art**

The art in biotechnology, as relates to the association of diseases with particular genes, is highly unpredictable. The claimed sequences currently appear to represent orphan genes, since no matches were identified in a sequence search. Regarding such Orphan genes, Dujon (Trends in Genetics (1996) 12(7):263-270) notes that the most striking result of yeast sequencing is that "a significant proportion of yeast genes are orphans of unpredictable function (abstract)". Dujon further states "We have no clue to which direction to search and, even worse, when considering the experiments that could be done on orphans, we rapidly find ourselves intellectually embedded in the schemes of the past (page 2169, column 2)." Thus, it is extremely unpredictable what to do with an orphan gene such as SEQ ID NOs: 1-10 in the absence of any defined utility.

Further, as noted above, the sequences are not even species or chromosome specific, based upon the sequence search. As the attached search of SEQ ID NO: 3 in Genbank demonstrates, result 3 shows an 90% match (local similarity) to a sequence in chromosome 14 of humans while result 13 shows an 89.5% match (local similarity) to human chromosome 8. Further, the remaining results show similarity to Pseudomonas, Lotus, Rats and Rice with equivalent levels of local similarity. Similar results exist for the other 9 probes. So the sequences claimed lack 100% specificity to any particular organism in Genbank, and the specification lacks any discussion of the target for these oligonucleotides. Consequently, there is no specific target for any of the claimed sequences. In the absence of any target, it is entirely unpredictable how these sequences would function even in some sort of genomic analysis method. SEQ ID NO:

3, for example, would crosshybridize to both chromosomes 8 and 14 and would not give significant information regarding the presence or absence of any particular human, animal or plant sequence in a sample, since the sequence would hybridize about equally well to human, rat, rice and lotus, among other species.

### **Quantity of Experimentation**

An immense amount of experimentation would be required in order to define whether any of these nucleic acids are associated with any particular disease state or other specific and substantial use. For example, in order to acquire statistically significant evidence of an association with a disease or other utility, one of the possible targets such as human patients, experimental rats, or rice and lotus plants in each of the many hundreds of different possible disease states would need to be subjected to collection of samples for analysis of their DNA, followed by analysis and the inventive efforts of determining if any association exists. This is a very large quantity of experimentation.

### **Determination**

In view of the unpredictable nature of the invention, the absence of any guidance in the specification for a substantial and specific use, the absence of any working examples in the specification, the negative teachings in the prior art, the extreme unpredictability of the invention, and the large amount of experimentation necessary balanced against the high level of skill in the art and the relatively narrow breadth of the

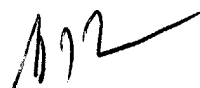
claims, it is concluded that undue experimentation would be required to use this invention as claimed.

### ***Conclusion***

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Jeffrey Fredman whose telephone number is currently 703-308-6568. In mid January, 2004, when TC 1600 relocates to the new USPTO facility in Alexandria, the examiner's phone number will become 571-272-0742. The examiner can normally be reached on 6:30-4:00.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Gary Benzion can be reached on 703-308-1119. The supervisor's new telephone number in mid January will be 571-272-0782. The fax phone number for the organization where this application or proceeding is assigned is currently 703-872-9306.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the receptionist whose telephone number is 703-308-0196.

  
Jeffrey Fredman  
Primary Examiner  
Art Unit 1634